

Iñigo Olalde

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

4,193
citations

218677

26
h-index

243625

44
g-index

59
all docs

59
docs citations

59
times ranked

4854
citing authors

#	ARTICLE	IF	CITATIONS
1	The Beaker phenomenon and the genomic transformation of northwest Europe. <i>Nature</i> , 2018, 555, 190-196.	27.8	503
2	The genomic history of southeastern Europe. <i>Nature</i> , 2018, 555, 197-203.	27.8	479
3	The formation of human populations in South and Central Asia. <i>Science</i> , 2019, 365, .	12.6	383
4	The genomic history of the Iberian Peninsula over the past 8000 years. <i>Science</i> , 2019, 363, 1230-1234.	12.6	340
5	Derived immune and ancestral pigmentation alleles in a 7,000-year-old Mesolithic European. <i>Nature</i> , 2014, 507, 225-228.	27.8	328
6	Reconstructing Prehistoric African Population Structure. <i>Cell</i> , 2017, 171, 59-71.e21.	28.9	308
7	Genomic Affinities of Two 7,000-Year-Old Iberian Hunter-Gatherers. <i>Current Biology</i> , 2012, 22, 1494-1499.	3.9	160
8	Ancient genomes indicate population replacement in Early Neolithic Britain. <i>Nature Ecology and Evolution</i> , 2019, 3, 765-771.	7.8	156
9	A Common Genetic Origin for Early Farmers from Mediterranean Cardial and Central European LBK Cultures. <i>Molecular Biology and Evolution</i> , 2015, 32, msv181.	8.9	155
10	Palaeo-Eskimo genetic ancestry and the peopling of Chukotka and North America. <i>Nature</i> , 2019, 570, 236-240.	27.8	118
11	Ancient DNA reveals a multistep spread of the first herders into sub-Saharan Africa. <i>Science</i> , 2019, 365, .	12.6	96
12	The spread of steppe and Iranian-related ancestry in the islands of the western Mediterranean. <i>Nature Ecology and Evolution</i> , 2020, 4, 334-345.	7.8	95
13	Ancient West African foragers in the context of African population history. <i>Nature</i> , 2020, 577, 665-670.	27.8	86
14	Large-scale migration into Britain during the Middle to Late Bronze Age. <i>Nature</i> , 2022, 601, 588-594.	27.8	86
15	Limiting replication stress during somatic cell reprogramming reduces genomic instability in induced pluripotent stem cells. <i>Nature Communications</i> , 2015, 6, 8036.	12.8	84
16	A genetic history of the pre-contact Caribbean. <i>Nature</i> , 2021, 590, 103-110.	27.8	67
17	A high-resolution picture of kinship practices in an Early Neolithic tomb. <i>Nature</i> , 2022, 601, 584-587.	27.8	65
18	Ancient DNA and deep population structure in sub-Saharan African foragers. <i>Nature</i> , 2022, 603, 290-296.	27.8	51

#	ARTICLE	IF	CITATIONS
19	On the path to extinction: Inbreeding and admixture in a declining grey wolf population. <i>Molecular Ecology</i> , 2018, 27, 3599-3612.	3.9	46
20	Mitochondrial DNA from the eradicated European <i>Plasmodium vivax</i> and <i>P. falciparum</i> from 70-year-old slides from the Ebro Delta in Spain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 11495-11500.	7.1	41
21	Genome data from a sixteenth century pig illuminate modern breed relationships. <i>Heredity</i> , 2015, 114, 175-184.	2.6	39
22	Genomic transformation and social organization during the Copper Age–Bronze Age transition in southern Iberia. <i>Science Advances</i> , 2021, 7, eabi7038.	10.3	39
23	<i>Plasmodium vivax</i> Malaria Viewed through the Lens of an Eradicated European Strain. <i>Molecular Biology and Evolution</i> , 2020, 37, 773-785.	8.9	38
24	Kinship and social organization in Copper Age Europe. A cross-disciplinary analysis of archaeology, DNA, isotopes, and anthropology from two Bell Beaker cemeteries. <i>PLoS ONE</i> , 2020, 15, e0241278.	2.5	35
25	Evolutionary and functional impact of common polymorphic inversions in the human genome. <i>Nature Communications</i> , 2019, 10, 4222.	12.8	34
26	Malaria was a weak selective force in ancient Europeans. <i>Scientific Reports</i> , 2017, 7, 1377.	3.3	32
27	Stone Age <i>Yersinia pestis</i> genomes shed light on the early evolution, diversity, and ecology of plague. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2116722119.	7.1	31
28	Mitochondrial DNA from El Mirador Cave (Atapuerca, Spain) Reveals the Heterogeneity of Chalcolithic Populations. <i>PLoS ONE</i> , 2014, 9, e105105.	2.5	28
29	Ancient DNA reveals monozygotic newborn twins from the Upper Palaeolithic. <i>Communications Biology</i> , 2020, 3, 650.	4.4	25
30	Population connectivity buffers genetic diversity loss in a seabird. <i>Frontiers in Zoology</i> , 2013, 10, 28.	2.0	20
31	Analysis of structural diversity in wolf-like canids reveals post-domestication variants. <i>BMC Genomics</i> , 2014, 15, 465.	2.8	16
32	Genomic analysis of the blood attributed to Louis XVI (1754–1793), king of France. <i>Scientific Reports</i> , 2015, 4, 4666.	3.3	16
33	Latest trends in archaeogenetic research of west Eurasians. <i>Current Opinion in Genetics and Development</i> , 2020, 62, 36-43.	3.3	16
34	Genetic comparison of the head of Henri IV and the presumptive blood from Louis XVI (both Kings of France). <i>PLoS ONE</i> , 2017, 12, e0170000.	2.2	13
35	Ancient DNA sheds light on the ancestry of pre-hispanic Canarian pigs. <i>Genetics Selection Evolution</i> , 2015, 47, 40.	3.0	13
36	Social stratification without genetic differentiation at the site of Kulubnarti in Christian Period Nubia. <i>Nature Communications</i> , 2021, 12, 7283.	12.8	13

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37	Genome-wide analysis of nearly all the victims of a 6200 year old massacre. PLoS ONE, 2021, 16, e0247332.	2.5	11
38	At the beginnings of the funerary Megalithism in Iberia at Campo de Hockey necropolis. Scientific Reports, 2022, 12, .	3.3	8
39	Combining ancient DNA and radiocarbon dating data to increase chronological accuracy. Journal of Archaeological Science, 2021, 133, 105452.	2.4	7
40	Modern humans' paleogenomics and the new evidences on the European prehistory. Science and Technology of Archaeological Research, 2015, 1, 1-9.	2.4	5
41	â€˜Avaâ€™: a Beaker-associated woman from a cist at Achavanich, Highland, and the story of her (re-)discovery and subsequent study. Proceedings of the Society of Antiquaries of Scotland, 0, 147, 73-118.	0.0	3
42	Salmonella enterica from a soldier from the 1652 siege of Barcelona (Spain) supports historical transatlantic epidemic contacts. IScience, 2021, 24, 103021.	4.1	2
43	Genomic Analysis of 18th-Century Kazakh Individuals and Their Oral Microbiome. Biology, 2021, 10, 1324.	2.8	2